

## RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/731,457A

Source: 1646

Date Processed by STIC: 5-17-01

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST 25. Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

**<http://www.uspto.gov/web/offices/pac/checker>**

1646

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/731,457A

DATE: 05/17/2001

TIME: 10:49:40

Input Set : A:\es.txt

Output Set: N:\CRF3\05172001\I731457A.raw

Does Not Comply  
Corrected Diskette Needed

4 <110> APPLICANT: Ian Popoff  
5 Jacqueline Wyatt  
7 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF DAMAGE-SPECIFIC DNA BINDING PROTEIN  
1, P127  
8 EXPRESSION  
10 <130> FILE REFERENCE: RTS-0182  
12 <140> CURRENT APPLICATION NUMBER: US/09/731,457A  
13 <141> CURRENT FILING DATE: 2000-12-06  
15 <160> NUMBER OF SEQ ID NOS: 87

## ERRORED SEQUENCES

436 <210> SEQ ID NO: 9  
437 <211> LENGTH: (20)  
438 <212> TYPE: DNA  
439 <213> ORGANISM: Artificial Sequence  
441 <220> FEATURE:  
443 <223> OTHER INFORMATION: PCR Probe  
445 <400> SEQUENCE: 9  
E--> 446 caagcttccc gttctcagcc

*Number of nucleic bases differ:**- 20 shown**- 21 instead*

(21)

VERIFICATION SUMMARY

DATE: 05/17/2001

PATENT APPLICATION: US/09/731,457A

TIME: 10:49:41

Input Set : A:\es.txt

Output Set: N:\CRF3\05172001\I731457A.raw

L:446 M:254 E: No. of Bases conflict, LENGTH:Input:21 Counted:20 SEQ:9